Validation for the determination of MDA in human plasma: Intra-assay precision and accuracy of quality control sample data in water

TABLE 1a

Validation for the determination of MDA in human plasma: Intra-assay precision and accuracy of quality control sample data in plasma

Replicate	QC 0.4 µmol/1. (LLOQ QC) Observed concentration (µmol/L)	QC 1 µmol/L (LoQC) Observed concentration (µmol/L)	QC 4 µmol/L (MeQC) Observed concentration (µmol/L)	QC 8 µmol/L (HiQC) Observed concentration (µmol/L)
1	0.39	0.92	3.78	7.65
2	0.43	0.95	3.91	7.50
3	0.44	1.05	4.03	7.87
4	0.45	1.05	3.72	8.37
5	NR*	NR*	4.44	8.13
6	0.68*	1.62*	4.28	8.37
Mean (μmol/L)	0.43	0.99	4.03_	7.98
SD (a-1)	0.026	0.068	0.284	0.368
Precision (%)	6.0	6.9	7.0	4.6
Accuracy (%)	107.5	99.0	100.8	99.8

	QC 1 μmol/L (LoQC)	QC 4 μmol/L (MeQC)	QC 8 μmol/L (HiQC)	
Replicate	Observed concentration (µmol/L)	Observed concentration (µmol/L)	Observed concentration (µmol/L)	
	•			
1	0.95	3.79	7.34	
2	1.00	3.87	7.34	
3	0.97	4.05	7.58	
4	0.92	3.92	7.63	
5	0.91	3.88	7.57	
6	0.90	3.88	7.82	
Mean (µmol/L)	0.94	3.90	7.55	
SD (n-1)	0.039	0.086	0.184	
Precision (%)	4.1	2.2	2.4	
Accuracy (%)	94.0	97.5	94.4	

NR = No result

^{*} two samples possibly transferred to same tube in error, result not included in statistics

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Validation for the determination of MDA in human plasma:

Inter-assay precision and accuracy of quality control sample data in water

TABLE 2

TABLE 2a

Validation for the determination of MDA in human plasma: Inter-assay precision and accuracy of quality control sample data in water

Batch	Replicate	QC 0.4 µmol/I. (LLOQ QC) Observed concentration (µmol/L)	QC 1 µmol/L (LoQC) Observed concentration (µmol/L)	QC 4 µmol/L (MeQC) Observed concentration (µmol/L)	QC 8 µmol/l. (HiQC) Observed concentration (µmol/L)	Batch	Replicate	QC 1 µmol/L (LoQC) Observed concentration (µmol/L)	QC 4 µmol/L (MeQC) Observed concentration (µmol/L)	QC 8 µmol/L (HiQC) Observed concentration (µmol/L)
PVAL01RI	1	0.39	0.92	3.78	7.65	PVAL01RI	1	0.95	3.79	7.34
	2	0.43	0.95	3.91	7.50		2	1.00	3.87	7.34
	3	0.44	1.05	4.03	7.87		3	0.97	4.05	7.58
	4	0.45	1.05	3.72	8.37		4	0.92	3.92	7.63
	5	NR*	NR*	4.44	8.13		5	0.91	3.88	7.57
	6	0.68*	1.62*	4.28	8.37		6	0.90	3.88	7.82
PVAL021	1	0.36	0.91	3.70	7.33	PVAL021	1	0.84	3.56	7.02
	2	0.39	0.96	3.89	7.65		2	0.92	3.78	7.42
	3	0.34	1.03	3.98	7.99		3	0.94	3.83	7.55
	4	0.41	1.04	4.12	8.21		4	0.99	3.80	7.74
	5	0.42	1.05	4.14	8.14		5	0.90	3.94	7.75
	6	0.42	1.04	4.35	8.78		6	0.89	4.11	8.10

NR = No result

^{*} two samples possibly transferred to same tube in error, result not included in statistics

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TABLE 2 (continued)

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Validation for the determination of MDA in human plasma: Inter-assay precision and accuracy of quality control sample data in water

TABLE 2a (continued)

Validation for the determination of MDA in human plasma: Inter-assay precision and accuracy of quality control sample data in plasma

		QC 0.4 µmol/L (LLOQ QC)	QC 1 μmol/L (LoQC)	QC 4 μmol/L (MeQC)	QC 8 µmol/L (HiQC)	D . 1		QC 1 µmot/L (LoQC)	QC 4 μmol/L (MeQC)	QC 8 µmol/L (HiQC)
Batch	Replicate	Observed concentration (µmol/L)	Observed concentration (µmol/L)	Observed concentration (µmol/L)	Observed concentration (µmol/L)	Batch	Replicate	Observed concentration (µmol/L)	Observed concentration (µmol/L)	Observed concentration (µmol/L)
PVAL03	ļ	•	0.90	3.71	7.34	PVAL03	1	0.99	3.59	6.85
	2	-	0.94	3.78	7.33		2	0.95	3.59	7.07
	3	-	0.97	3.88	7.96		3	0.92	3.90	7.38
	4	-	1.04	4.15	8.26		4	1.03	3.88	7.71
	5	-	1.06	4.27	8.50		5	0.99	4.15	7.71
	6	•	1.09	4.28	8.38		6	0.99	3.95	7.92
PVAL05	1	-	0.88	3.74	7.55	PVAL05	1	0.85	3.54	6,89
	2	-	0.97	3.90	7.91		2	0.92	3.68	7.15
	3	•	0.93	3.95	8.04		3	0.96	3.74	7.23
	4	-	0.97	4.07	8.11		4	0.94	3.82	7.56
	5	•	0.99	4.25	8.63		5	1.00	4.04	7.91
	6	-	1.03	4.36	8.64		6	1.10	3.94	7.68
Mean (µmo	VL)	0.41	0.99	4.03	8.03	Mean (μmol/	L)	0.95	3.84	7.50
Standard de	viation (n-1)	0.035	0.061	0.233	0.432	Standard dev	iation (n-I)	0.058	0.167	0.334
Precision (%	ú)	8.5	6.2	5.8	5.4	Precision (%))	6.1	4.3	4.5
Ассигасу (⁹ /	(i)	102.5	99.0	100.8	100.4	Accuracy (%)	95.0	96,0	93.8
n	·	10	22	24	24	ຄ		24	24	24

TABLE 3

Validation for the determination of MDA in human plasma: Inter-assay accuracy and precision of calibration standard data

			I	Back-calculate	ed concentra	itions (µmol/	'L)				Curve parameters	
Batch				Calibra	ation level (p	umol/L)				Gradient (m)	Intercept (c)	Coefficient of determination (r ²)
	0.4 0.6 1 2 3 5 7.5 9 10		10									
PVAL01RI	0.42	0.65	0.88	1.66 РГ	3.03	4.63	7.71	8.89	10.28	8.55660E 04	1.02740E 04	0.99742
PVAL02I	0.46	0.60	0.95	1.87	3.06	4.34	7.65	9.71	9.85	5.64649E 04	9.73083E 03	0.99147
PVAL03	0.35	0.67	1.04	1.88	3.10	4.86	7.74	9.74	9.12	6,66945E 04	4.47859E 03	0.98721
PVAL05	0.35	0.64	1.05	1.97	3.17	5.01	7.53	7.79	11.00	6.29813E 04	7.25626E 03	0.97809
Mean (µmol/L)	0.40	0.64	0.98	1.91	3.09	4.71	7.66	9.03	10.06			<u> </u>
Standard deviation (n-1)	0.054	0.029	0.080	0.055	0.061	0.292	0.093	0.917	0.787	-	-	-
Precision (%)	13.5	4.5	8.2	2.9	2.0	6.2	1.2	10.2	7.8	-	•	•
Accuracy (%)	100.0	106.7	98.0	95.5	103.0	94.2	102.1	100.3	100.6		-	-

PF - Poor fit, calibration standard outside acceptance criteria

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TABLE 4

Validation for the determination of MDA in human plasma:

Individual unspiked plasma samples

(endogenous levels)

Matrix sample number	Back calculated concentration (umol/L)	Mean back calculated concentrations (unol/L) (CV%)
01H408	0.57	0.6
01H409	0.62	(10.0)
01H410	0.61	
01H411	0.67	
01H412	0.57	
01H319	0.48	
01H182	0,66	
01H183	0.58	
01H184	0.66	
01H318	0.70	

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TABLE 5

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Validation for the determination of MDA in human plasma: Recovery of MDA from human plasma

Quality control level (µmol/L)	Replicate	Extracted QCs. Conc (µmol/L)	Mean peak area (CV%)	Reference QCs. Cone (µmol/L)	Mean peak area (CV%)	Recovery (%)
1	i	0.99	1.0	0.90	1.0	100.0
	2	0.95	(4.0)	0.94	(7.0)	
	3	0.92		0.97		
	4	1.03		1,04		
	5	0.99		1,06		
	6	0.99		1.09		
4	1	3.59	3.8	3.71	4.0	94.7
	2	3.59	(5.7)	3.78	(6.3)	
	3	3.90		3,88		
	4	3,88		4.15		
	5	4.15		4.27		
	6	3.95		4.28		
8	1	6.85	7.4	7.34	8.0	92.5
	2	7.07	(5.6)	7.33	(6.5)	
	3	7.38		7.96		
	4	7.71		8.26		
	5	7,71		8,50		
	6	7.92		8.38		
fean recovery (%)	<u></u>					95.7

Validation for the determination of MDA in human plasma: Room temperature stability of MDA in human plasma (24 hours)

TABLE 6

Quality control level (μmol/L)		Baseline	QC samples	QC samples stored extra		
	Replicate	Observed concentration (µmol/L)	Mean (μmol/L) (CV%)	Observed concentration (µmol/L)	Mean (μmol/L) (CV%)	Difference from baseline (%)
1	1	0.99	0.98	0.87	0.89	-9.2
	2	0.95	(3.9)	0.84	(4.4)	
	3	0.92		0.95		
	4	1.03		0.89		
	5	0.99		0.89		
	6	0.99		0.88		
8	1	6.85	7.44	6.02	6.17	-17.1
	2	7.07	(5.6)	6.00	(2.3)	
	3	7.38		6.11		
	4	7.71		6.22		
	5	7.71		6.28		
	6	7.92		6.36		

Validation for the determination of MDA in human plasma; Room temperature stability of MDA in human plasma (18 hours)

TABLE 6a

Quality control level (µmol/L)		Baseline	QC samples	•	for 18 hours prior to action	
	Replicate	Observed concentration (µmol/L)	Mean (μmol/L) (CV%)	Observed concentration (µmol/L)	Mean (μmol/L) (CV%)	Difference from baseline (%)
1	1	0.85	0.96	0.87	0.90	-6.2
	2	0.92	(8.8)	0.88	(1.8)	
	3	0.96		0.91		
	4	0.94		0.92		
	5	1.00		0.91		
	6	1.10		0.92		
8	1	6.89	7.40	6.24	6.58	-11.1
	2	7.15	(5.1)	6.24	(4.9)	
	3	7.23		6.49		
	4	7.56		6.66		
	5	7.91		6.77		
	6	7.68		7.10		

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Validation for the determination of MDA in human plasma:
Room temperature stability of MDA in human plasma
(12 hours)

TABLE 6b

Quality control level (µmol/L)		Baseline	QC samples	•	for 12 hours prior to action	
	Replicate	Observed concentration (µmoi/L)	Mean (μmol/L) (CV%)	Observed concentration (µmol/L)	Mean (μmol/L) (CV%)	Difference from baseline (%)
1	1	0.85	0.96	0.86	0.90	-6.2
	2	0.92	(8.8)	0.85	(6.3)	
	3	0.96		0.87		
	4	0.94		0.89		
	5	1.00		0.97		
	6	1.10		0.97		
8	1	6.89	7.40	6.47	6.83	-7 .7
	2	7.15	(5.1)	6.58	(4.3)	
	3	7.23		6.59		
	4	7.56		7.10		
	5	7.91		7.03		
	6	7.68		7.19		

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Validation for the determination of MDA in human plasma: Freeze/thaw stability of MDA in human plasma

TABLE 7

Quality control Re level (µmol/L)	·	Baseline	QC samples		cted to 3 additional prior to extraction	
	Replicate	Observed concentration (µmol/L)	Mean (μmol/L) (CV%)	Observed concentration (µmol/L)	Mean (μmol/L) (CV%)	Difference from baseline (%)
1	1	0.84	0,91	0.81	0.88	-3.3
	2	0.92	(5.5)	0.88	(4.2)	
	3	0.94		0.87		
	4	0.99		0.87		
	5	0.90		0.92		
	6	0.89		0.90		
8	1	7.02	7.60	6.66	7.39	-2.8
	2	7.42	(4.8)	7.22	(5.7)	
	3	7.55		7.36		
	4	7.74		7.53		
	5	7.75		7.71		
	6	8.10		7.85		

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TABLE 8

Validation for the determination of MDA in human plasma: Stability of extracted QC samples from plasma, stored refrigerated for 3 days

TABLE 8a

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Validation for the determination of MDA in human plasma: Stability of extracted QC samples from water, stored refrigerated for 3 days

		Stored extract	Stored extracted QC samples					Stored extracted QC samples					
Quality control level (µmol/L)	Replicate	Observed concentration (µmol/L)	Mean (µmol/L) (CV%)	Difference from nominal concentration (%)	Quality control level (µmol/L)	Replicate	Observed concentration (µmol/L)	Mean (µmol/L) (CV%)	Difference from nominal concentration (%)				
1	1		#DIV/0!	#DIV/0!	1	1		#DIV/0!	#DIV/0!				
	2		#DIV/0!			2		#DIV/0!					
	3					3							
	4					4							
	5		•			5							
	6					6							
4	1		#DIV/0!	#DIV/0!	4	1		#DIV/0!	#DIV/0!				
	2		#DIV/0!			2		#DIV/0!					
	3					3							
	4					4							
	5					5							
	6					6							
8	1		#DIV/0!	#DIV/0!	8	1		#DIV/0!	#DIV/0!				
	2		#DIV/0!			2		#DIV/0!					
	3					3							
	4					4							
	5					5							
	6					6							

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Validation for the determination of MDA in human plasma: Stability of stock solutions stored at nominal 4°C for 29 days

TABLE 9

Date of stock solution preparation	Replicate	Peak absorbance of stock solution	Mean peak absorbance value	% Difference of stored solution from new solution
13/08/01 (new)	1		#DIV/0!	#DIV/0!
	2			
	3			
16/07/01 (stored)	1		#DIV/0!	
	2			
	3			

TABLE 3

Validation for the determination of MDA in human plasma:

Response function batch accuracy and precision of calibration standard data

	Rack-calculated concentrations (umol/L)									Curve parameters			
Matrix		Calibration level (µmol/L)								Gradient (m)	Intercept (c)	Coefficient of determination (r ²)	
	0.4	0.6	1	2	3	5	7.5	9	10			· · · · · · · · · · · · · · · · · · ·	
Water	0.44	0.62	0.62 PF	1.86	2.81	4.75	7.42	9.19	10.41	5.60778E 04	5.970631E 03	0.99688	
0111408	0.43	0.62	0.98	1.89	2.92	4.79	7.36	9.17	10.34	5.39809E 04	2.22627E 04	0.99806	
0111409	0.40	0.65	0.99	1.91	2.94	4.81	7.32	9.04	10.45	5.38390E 04	2.85398E 04	0.99756	
0111410	0.42	0.66	0.89	1.98	2.92	4.77	7.36	9.14	10.36	5.38928E 04	3.72385E 04	0.99789	
0111411	0.44	0.62	0.94	1.86	2.94	5.01	7.36	8.96	10.37	5.37681E 04	3.54146E 04	0.99836	
0111412	0.43	0.61	0.94	2.00	2.97	4.91	7.27	8.98	10.40	5.40267E 04	2.40573E 04	0.99803	
01II3 19	0.43	0.60	0.97	1.90	3.93 PF	4.97	7.54	8.97	10.12	5.37025E 04	2.28820E 04	0.99974	
Mean (µmol/L)	0.43	0.63	0.95	1.91	2.92	4.86	7.38	9.06	10.35	54184.0	-	-	
Standard deviation (n-1)	0.014	0.021	0.037	0.055	0.055	0.104	0.086	0.100	0.108	842.72	-	•	
Precision (%)	3.3	3.3	3.9	2.9	1.9	2.1	1.2	1.1	1.0	1.6	-	•	
Accuracy (%)	107.5	105.0	95.0	95.5	97.3	97.2	98.4	100.7	103.5	•	-	-	

PI = Poor fit, calibration standard outside acceptance criteria